

SEQUENCE LISTING

<110> Korneluk, Robert G.
MacKenzie, Alexander E.
Baird, Stephen
Liston, Peter

<120> MAMMALIAN IAP GENE FAMILY, PRIMERS,
PROBES, AND DETECTION METHODS

<130> 07891/003006

<150> US 09/011,356
<151> 1998-02-04

<150> PCT/IB96/01022
<151> 1996-08-05

<150> US 08/576,956
<151> 1995-12-22

<150> US 08/511,485
<151> 1995-08-04

<160> 45

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<213> Artificial Sequence

<220>
<223> Synthetic based on Homo sapiens, Mus musculus,
Drosophila melanogaster, Cydia pomonella, and
Orgyia pseudotsugata

<221> VARIANT
<222> (2)...(45)
<223> Xaa at positions 2, 3, 4, 5, 6, 7, 9, 10, 11, 17,
18, 19, 20, 21, 23, 25, 30, 31, 32, 34, 35, 38,
39, 40, 41, 42, and 45 may be any amino acid.

<221> VARIANT
<222> (8)...(8)
<223> Xaa at position 8 is Glu or Asp.

<221> VARIANT
<222> (14)...(14)
<223> Xaa at position 14 is Val or Ile.

<221> VARIANT
<222> (22)...(22)
<223> Xaa at position 22 is Val or Ile.

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 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Phe Xaa Pro Cys Gly His Xaa Xaa Xaa
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 Cys Xaa Xaa Cys Ala Xaa Xaa Xaa Xaa Xaa Cys Pro Xaa Cys
 35 40 45

<210> 2
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 <213> Artificial Sequence

<220>
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 <222> (1)...(66)
 <223> Xaa at positions 1, 2, 3, 6, 9, 10, 14, 15, 18,
 19, 20, 21, 24, 30, 32, 33, 35, 37, 40, 42, 43,
 44, 45, 46, 47, 49, 50, 51, 53, 54, 55, 56, 57,
 59, 60, 61, 62, 64 and 66 may be any amino acid.

<221> VARIANT
 <222> (13)...(17)
 <223> Xaa at positions 13, 16 and 17 may be any amino
 acid or may be absent.

<223> Synthetic based on Homo sapiens, Mus musculus,
 Drosophila melanogaster, Cydia pomonella, and
 Orgyia pseudotsugata

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 20 25 30
 Xaa Asp Xaa Val Xaa Cys Phe Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Trp
 35 40 45
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 <212> DNA
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<220>
 <221> variation
 <222> (2540)...(2540)
 <223> N may be any nucleotide

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 gcagggtttc ttatactgg tgaaggagat accgtgcggt gctttagttg tcatgcagct 240
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 <212> PRT
 <213> Homo sapiens

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 35 40 45
 Arg Ala Gly Phe Leu Tyr Thr Gly Glu Gly Asp Thr Val Arg Cys Phe
 50 55 60
 Ser Cys His Ala Ala Val Asp Arg Trp Gln Tyr Gly Asp Ser Ala Val
 65 70 75 80
 Gly Arg His Arg Lys Val Ser Pro Asn Cys Arg Phe Ile Asn Gly Phe
 85 90 95
 Tyr Leu Glu Asn Ser Ala Thr Gln Ser Thr Asn Ser Gly Ile Gln Asn
 100 105 110

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Gln	Val	Val	Asp	Ile	Ser	Asp	Thr	Ile	Tyr	Pro	Arg	Asn	Pro	Ala	Met	
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Tyr	Cys	Glu	Glu	Ala	Arg	Leu	Lys	Ser	Phe	Gln	Asn	Trp	Pro	Asp	Tyr	
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Ala	His	Leu	Thr	Pro	Arg	Glu	Leu	Ala	Ser	Ala	Gly	Leu	Tyr	Tyr	Thr	
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 Ser Leu Ala Arg Ala Gly Phe Tyr Tyr Thr Gly Val Asn Asp Lys Val
 50 55 60
 Lys Cys Phe Cys Cys Gly Leu Met Leu Asp Asn Trp Lys Arg Gly Asp
 65 70 75 80
 Ser Pro Thr Glu Lys His Lys Lys Leu Tyr Pro Ser Cys Arg Phe Val
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 Gln Ser Leu Asn Ser Val Asn Asn Leu Glu Ala Thr Ser Gln Pro Thr
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 Phe Pro Ser Ser Val Thr His Ser Thr His Ser Leu Leu Pro Gly Thr
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 Glu Asn Ser Gly Tyr Phe Arg Gly Ser Tyr Ser Asn Ser Pro Ser Asn
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 Ser Tyr Pro Cys Pro Met Asn Asn Glu Asn Ala Arg Leu Leu Thr Phe
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 Gln Thr Trp Pro Leu Thr Phe Leu Ser Pro Thr Asp Leu Ala Arg Ala
 180 185 190
 Gly Phe Tyr Tyr Ile Gly Pro Gly Asp Arg Val Ala Cys Phe Ala Cys
 195 200 205
 Gly Gly Lys Leu Ser Asn Trp Glu Pro Lys Asp Asn Ala Met Ser Glu
 210 215 220
 His Leu Arg His Phe Pro Lys Cys Pro Phe Ile Glu Asn Gln Leu Gln
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 Asp Thr Ser Arg Tyr Thr Val Ser Asn Leu Ser Met Gln Thr His Ala
 245 250 255
 Ala Arg Phe Lys Thr Phe Phe Asn Trp Pro Ser Ser Val Leu Val Asn
 260 265 270
 Pro Glu Gln Leu Ala Ser Ala Gly Phe Tyr Tyr Val Gly Asn Ser Asp
 275 280 285
 Asp Val Lys Cys Phe Cys Cys Asp Gly Gly Leu Arg Cys Trp Glu Ser
 290 295 300
 Gly Asp Asp Pro Trp Val Gln His Ala Lys Trp Phe Pro Arg Cys Glu
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 Tyr Leu Ile Arg Ile Lys Gly Gln Glu Phe Ile Arg Gln Val Gln Ala
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 Ser Tyr Pro His Leu Leu Glu Gln Leu Leu Ser Thr Ser Asp Ser Pro
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 Gly Asp Glu Asn Ala Glu Ser Ser Ile Ile His Leu Glu Pro Gly Glu
 355 360 365
 Asp His Ser Glu Asp Ala Ile Met Met Asn Thr Pro Val Ile Asn Ala
 370 375 380
 Ala Val Glu Met Gly Phe Ser Arg Ser Leu Val Lys Gln Thr Val Gln
 385 390 395 400

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Val	Ile	Lys	Gln	Lys	Thr	Gln	Thr	Ser	Leu	Gln	Ala	Arg	Glu	Leu	Ile
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Asp	Thr	Ile	Leu	Val	Lys	Gly	Asn	Ile	Ala	Ala	Thr	Val	Phe	Arg	Asn
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Met	Asp	Lys	Glu	Val	Ser	Ile	Val	Phe	Ile	Pro	Cys	Gly	His	Leu	Val
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Val	Cys	Lys	Asp	Cys	Ala	Pro	Ser	Leu	Arg	Lys	Cys	Pro	Ile	Cys	Arg
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<211> 2580

<212> DNA

<213> Homo sapiens

<220>

<221> variation

<222> (2412)...(2412)

<223> N may be any nucleotide

<400> 7

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catgtgaaga	aatttcatgt	gaatgtttta	gctatcaa	agtactgtca	cctactcatg	240
cacaaaactg	cctcccaaag	acttttccca	ggtccctcgt	atcaaaacat	taagagtata	300
atggaagata	gcacgatctt	gtcagattgg	acaaacagca	acaaacaaaa	aatgaagtat	360
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<210> 8
 <211> 618
 <212> PRT
 <213> Homo sapiens

<400> 8

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			20					25					30		
Asn	Ser	Asn	Lys	Gln	Lys	Met	Lys	Tyr	Asp	Phe	Ser	Cys	Glu	Leu	Tyr
		35					40					45			
Arg	Met	Ser	Thr	Tyr	Ser	Thr	Phe	Pro	Ala	Gly	Val	Pro	Val	Ser	Glu
	50					55					60				
Arg	Ser	Leu	Ala	Arg	Ala	Gly	Phe	Tyr	Tyr	Thr	Gly	Val	Asn	Asp	Lys
65					70					75					80
Val	Lys	Cys	Phe	Cys	Cys	Gly	Leu	Met	Leu	Asp	Asn	Trp	Lys	Leu	Gly
			85					90						95	
Asp	Ser	Pro	Ile	Gln	Lys	His	Lys	Gln	Leu	Tyr	Pro	Ser	Cys	Ser	Phe
			100					105					110		
Ile	Gln	Asn	Leu	Val	Ser	Ala	Ser	Leu	Gly	Ser	Thr	Ser	Lys	Asn	Thr
		115					120					125			
Ser	Pro	Met	Arg	Asn	Ser	Phe	Ala	His	Ser	Leu	Ser	Pro	Thr	Leu	Glu
		130				135					140				
His	Ser	Ser	Leu	Phe	Ser	Gly	Ser	Tyr	Ser	Ser	Leu	Pro	Pro	Asn	Pro
145					150					155					160
Leu	Asn	Ser	Arg	Ala	Val	Glu	Asp	Ile	Ser	Ser	Ser	Arg	Thr	Asn	Pro
			165					170						175	
Tyr	Ser	Tyr	Ala	Met	Ser	Thr	Glu	Glu	Ala	Arg	Phe	Leu	Thr	Tyr	His
			180					185					190		
Met	Trp	Pro	Leu	Thr	Phe	Leu	Ser	Pro	Ser	Glu	Leu	Ala	Arg	Ala	Gly
		195					200					205			
Phe	Tyr	Tyr	Ile	Gly	Pro	Gly	Asp	Arg	Val	Ala	Cys	Phe	Ala	Cys	Gly
	210					215					220				

Gly	Lys	Leu	Ser	Asn	Trp	Glu	Pro	Lys	Asp	Asp	Ala	Met	Ser	Glu	His	225	230	235	240
Arg	Arg	His	Phe	Pro	Asn	Cys	Pro	Phe	Leu	Glu	Asn	Ser	Leu	Glu	Thr	245	250	255	
Leu	Arg	Phe	Ser	Ile	Ser	Asn	Leu	Ser	Met	Gln	Thr	His	Ala	Ala	Arg	260	265	270	
Met	Arg	Thr	Phe	Met	Tyr	Trp	Pro	Ser	Ser	Val	Pro	Val	Gln	Pro	Glu	275	280	285	
Gln	Leu	Ala	Ser	Ala	Gly	Phe	Tyr	Tyr	Val	Gly	Arg	Asn	Asp	Asp	Val	290	295	300	
Lys	Cys	Phe	Gly	Cys	Asp	Gly	Gly	Leu	Arg	Cys	Trp	Glu	Ser	Gly	Asp	305	310	315	320
Asp	Pro	Trp	Val	Glu	His	Ala	Lys	Trp	Phe	Pro	Arg	Cys	Glu	Phe	Leu	325	330	335	
Ile	Arg	Met	Lys	Gly	Gln	Glu	Phe	Val	Asp	Glu	Ile	Gln	Gly	Arg	Tyr	340	345	350	
Pro	His	Leu	Leu	Glu	Gln	Leu	Leu	Ser	Thr	Ser	Asp	Thr	Thr	Gly	Glu	355	360	365	
Glu	Asn	Ala	Asp	Pro	Pro	Ile	His	Phe	Gly	Pro	Gly	Glu	Ser	Ser		370	375	380	
Ser	Glu	Asp	Ala	Val	Met	Met	Asn	Thr	Pro	Val	Val	Lys	Ser	Ala	Leu	385	390	395	400
Glu	Met	Gly	Phe	Asn	Arg	Asp	Leu	Val	Lys	Gln	Thr	Val	Leu	Ser	Lys	405	410	415	
Ile	Leu	Thr	Thr	Gly	Glu	Asn	Tyr	Lys	Thr	Val	Asn	Asp	Ile	Val	Ser	420	425	430	
Ala	Leu	Leu	Asn	Ala	Glu	Asp	Glu	Lys	Arg	Glu	Glu	Glu	Lys	Glu	Lys	435	440	445	
Gln	Ala	Glu	Glu	Met	Ala	Ser	Asp	Asp	Leu	Ser	Leu	Ile	Arg	Lys	Asn	450	455	460	
Arg	Met	Ala	Leu	Phe	Gln	Gln	Leu	Thr	Cys	Val	Leu	Pro	Ile	Leu	Asp	465	470	475	480
Asn	Leu	Leu	Lys	Ala	Asn	Val	Ile	Asn	Lys	Gln	Glu	His	Asp	Ile	Ile	485	490	495	
Lys	Gln	Lys	Thr	Gln	Ile	Pro	Leu	Gln	Ala	Arg	Glu	Leu	Ile	Asp	Thr	500	505	510	
Ile	Trp	Val	Lys	Gly	Asn	Ala	Ala	Ala	Asn	Ile	Phe	Lys	Asn	Cys	Leu	515	520	525	
Lys	Glu	Ile	Asp	Ser	Thr	Leu	Tyr	Lys	Asn	Leu	Phe	Val	Asp	Lys	Asn	530	535	540	
Met	Lys	Tyr	Ile	Pro	Thr	Glu	Asp	Val	Ser	Gly	Leu	Ser	Leu	Glu	Glu	545	550	555	560
Gln	Leu	Arg	Arg	Leu	Gln	Glu	Glu	Arg	Thr	Cys	Lys	Val	Cys	Met	Asp	565	570	575	
Lys	Glu	Val	Ser	Val	Val	Phe	Ile	Pro	Cys	Gly	His	Leu	Val	Val	Cys	580	585	590	
Gln	Glu	Cys	Ala	Pro	Ser	Leu	Arg	Lys	Cys	Pro	Ile	Cys	Arg	Gly	Ile	595	600	605	
Ile	Lys	Gly	Thr	Val	Arg	Thr	Phe	Leu	Ser							610	615		

<210> 9

<211> 2100

<212> DNA

<213> Mus musculus

<400> 9

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60

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gagaagatga	cttttaacag	ttttgaagga	actagaactt	ttgtacttgc	agacaccaat	180
aaggatgaag	aatttgtaga	agagtttaat	agattaaaaa	catttgctaa	cttcccaagt	240
agtagtcctg	tttcagcatc	aacattggcg	cgagctgggt	ttctttatac	cgggtgaagga	300
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gaatctgccc	aatgacttta	attggcttat	tgtaaacacg	gaaagaactg	ccccacgctg	2040
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<210> 10

<211> 496

<212> PRT

<213> Mus musculus

<400> 10

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Thr	Asn	Lys	Asp	Glu	Glu	Phe	Val	Glu	Glu	Phe	Asn	Arg	Leu	Lys	Thr
			20					25					30		
Phe	Ala	Asn	Phe	Pro	Ser	Ser	Ser	Pro	Val	Ser	Ala	Ser	Thr	Leu	Ala
		35					40				45				
Arg	Ala	Gly	Phe	Leu	Tyr	Thr	Gly	Glu	Gly	Asp	Thr	Val	Gln	Cys	Phe
		50				55					60				
Ser	Cys	His	Ala	Ala	Ile	Asp	Arg	Trp	Gln	Tyr	Gly	Asp	Ser	Ala	Val
65					70				75					80	
Gly	Arg	His	Arg	Arg	Ile	Ser	Pro	Asn	Cys	Arg	Phe	Ile	Asn	Gly	Phe
			85					90					95		
Tyr	Phe	Glu	Asn	Gly	Ala	Ala	Gln	Ser	Thr	Asn	Pro	Gly	Ile	Gln	Asn
			100					105					110		
Gly	Gln	Tyr	Lys	Ser	Glu	Asn	Cys	Val	Gly	Asn	Arg	Asn	Pro	Phe	Ala
		115				120						125			
Pro	Asp	Arg	Pro	Pro	Glu	Thr	His	Ala	Asp	Tyr	Leu	Leu	Arg	Thr	Gly

130	135	140
Gln Val Val Asp Ile Ser Asp Thr Ile Tyr Pro Arg Asn Pro Ala Met		
145	150	155
Cys Ser Glu Glu Ala Arg Leu Lys Ser Phe Gln Asn Trp Pro Asp Tyr		160
	165	170
Ala His Leu Thr Pro Arg Glu Leu Ala Ser Ala Gly Leu Tyr Tyr Thr		175
	180	185
Gly Ala Asp Asp Gln Val Gln Cys Phe Cys Cys Gly Gly Lys Leu Lys		190
	195	200
Asn Trp Glu Pro Cys Asp Arg Ala Trp Ser Glu His Arg Arg His Phe		205
	210	215
Pro Asn Cys Phe Phe Val Leu Gly Arg Asn Val Asn Val Arg Ser Glu		220
225	230	235
Ser Gly Val Ser Ser Asp Arg Asn Phe Pro Asn Ser Thr Asn Ser Pro		240
	245	250
Arg Asn Pro Ala Met Ala Glu Tyr Glu Ala Arg Ile Val Thr Phe Gly		255
	260	265
Thr Trp Ile Tyr Ser Val Asn Lys Glu Gln Leu Ala Arg Ala Gly Phe		270
	275	280
Tyr Ala Leu Gly Glu Gly Asp Lys Val Lys Cys Phe His Cys Gly Gly		285
	290	295
Gly Leu Thr Asp Trp Lys Pro Ser Glu Asp Pro Trp Asp Gln His Ala		300
305	310	315
Lys Cys Tyr Pro Gly Cys Lys Tyr Leu Leu Asp Glu Lys Gly Gln Glu		320
	325	330
Tyr Ile Asn Asn Ile His Leu Thr His Pro Leu Glu Glu Ser Leu Gly		335
	340	345
Arg Thr Ala Glu Lys Thr Pro Pro Leu Thr Lys Lys Ile Asp Asp Thr		350
	355	360
Ile Phe Gln Asn Pro Met Val Gln Glu Ala Ile Arg Met Gly Phe Ser		365
	370	375
Phe Lys Asp Leu Lys Lys Thr Met Glu Glu Lys Ile Gln Thr Ser Gly		380
385	390	395
Ser Ser Tyr Leu Ser Leu Glu Val Leu Ile Ala Asp Leu Val Ser Ala		400
	405	410
Gln Lys Asp Asn Thr Glu Asp Glu Ser Ser Gln Thr Ser Leu Gln Lys		415
	420	425
Asp Ile Ser Thr Glu Glu Gln Leu Arg Arg Leu Gln Glu Glu Lys Leu		430
	435	440
Ser Lys Ile Cys Met Asp Arg Asn Ile Ala Ile Val Phe Phe Pro Cys		445
	450	455
Gly His Leu Ala Thr Cys Lys Gln Cys Ala Glu Ala Val Asp Lys Cys		460
465	470	475
Pro Met Cys Tyr Thr Val Ile Thr Phe Asn Gln Lys Ile Phe Met Ser		480
	485	490
		495

<210> 11

<211> 67

<212> PRT

<213> Orgyia pseudotsugata

<400> 11

Lys Ala Ala Arg Leu Gly Thr Tyr Thr Asn Trp Pro Val Gln Phe Leu	
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Glu Pro Ser Arg Met Ala Ala Ser Gly Phe Tyr Tyr Leu Gly Arg Gly	
	10
	15
	20
Asp Glu Val Arg Cys Ala Phe Cys Lys Val Glu Ile Thr Asn Trp Val	
	25
	30
	35
	40
	45

Arg Gly Asp Asp Pro Glu Thr Asp His Lys Arg Trp Ala Pro Gln Cys
 50 55 60
 Pro Phe Val
 65

<210> 12
 <211> 275
 <212> PRT
 <213> *Cydia pomonella*

<400> 12
 Met Ser Asp Leu Arg Leu Glu Glu Val Arg Leu Asn Thr Phe Glu Lys
 1 5 10 15
 Trp Pro Val Ser Phe Leu Ser Pro Glu Thr Met Ala Lys Asn Gly Phe
 20 25 30
 Tyr Tyr Leu Gly Arg Ser Asp Glu Val Arg Cys Ala Phe Cys Lys Val
 35 40 45
 Glu Ile Met Arg Trp Lys Glu Gly Glu Asp Pro Ala Ala Asp His Lys
 50 55 60
 Lys Trp Ala Pro Gln Cys Pro Phe Val Lys Gly Ile Asp Val Cys Gly
 65 70 75 80
 Ser Ile Val Thr Thr Asn Asn Ile Gln Asn Thr Thr Thr His Asp Thr
 85 90 95
 Ile Ile Gly Pro Ala His Pro Lys Tyr Ala His Glu Ala Ala Arg Val
 100 105 110
 Lys Ser Phe His Asn Trp Pro Arg Cys Met Lys Gln Arg Pro Glu Gln
 115 120 125
 Met Ala Asp Ala Gly Phe Phe Tyr Thr Gly Tyr Gly Asp Asn Thr Lys
 130 135 140
 Cys Phe Tyr Cys Asp Gly Gly Leu Lys Asp Trp Glu Pro Glu Asp Val
 145 150 155 160
 Pro Trp Glu Gln His Val Arg Trp Phe Asp Arg Cys Ala Tyr Val Gln
 165 170 175
 Leu Val Lys Gly Arg Asp Tyr Val Gln Lys Val Ile Thr Glu Ala Cys
 180 185 190
 Val Leu Pro Gly Glu Asn Thr Thr Val Ser Thr Ala Ala Pro Val Ser
 195 200 205
 Glu Pro Ile Pro Glu Thr Lys Ile Glu Lys Glu Pro Gln Val Glu Asp
 210 215 220
 Ser Lys Leu Cys Lys Ile Cys Tyr Val Glu Glu Cys Ile Val Cys Phe
 225 230 235 240
 Val Pro Cys Gly His Val Val Ala Cys Ala Lys Cys Ala Leu Ser Val
 245 250 255
 Asp Lys Cys Pro Met Cys Arg Lys Ile Val Thr Ser Val Leu Lys Val
 260 265 270
 Tyr Phe Ser
 275

<210> 13
 <211> 498
 <212> PRT
 <213> *Drosophila melanogaster*

<400> 13
 Met Thr Glu Leu Gly Met Glu Leu Glu Ser Val Arg Leu Ala Thr Phe
 1 5 10 15
 Gly Glu Trp Pro Leu Asn Ala Pro Val Ser Ala Glu Asp Leu Val Ala

Leu Ser

<210> 14
<211> 67
<212> PRT
<213> *Cydia pomonella*

<400> 14
Glu Glu Val Arg Leu Asn Thr Phe Glu Lys Trp Pro Val Ser Phe Leu
1 5 10 15
Ser Pro Glu Thr Met Ala Lys Asn Gly Phe Tyr Tyr Leu Gly Arg Ser
20 25 30
Asp Glu Val Arg Cys Ala Phe Cys Lys Val Glu Ile Met Arg Trp Lys
35 40 45
Glu Gly Glu Asp Pro Ala Ala Asp His Lys Lys Trp Ala Pro Gln Cys
50 55 60
Pro Phe Val
65

<210> 15
<211> 67
<212> PRT
<213> *Drosophila melanogaster*

<400> 15
Glu Ala Asn Arg Leu Val Thr Phe Lys Asp Trp Pro Asn Pro Asn Ile
1 5 10 15
Thr Pro Gln Ala Leu Ala Lys Ala Gly Phe Tyr Tyr Leu Asn Arg Leu
20 25 30
Asp His Val Lys Cys Val Trp Cys Asn Gly Val Ile Ala Lys Trp Glu
35 40 45
Lys Asn Asp Asn Ala Phe Glu Glu His Lys Arg Phe Phe Pro Gln Cys
50 55 60
Pro Arg Val
65

<210> 16
<211> 68
<212> PRT
<213> *Mus musculus*

<400> 16
Glu Phe Asn Arg Leu Lys Thr Phe Ala Asn Phe Pro Ser Ser Ser Pro
1 5 10 15
Val Ser Ala Ser Thr Leu Ala Arg Ala Gly Phe Leu Tyr Thr Gly Glu
20 25 30
Gly Asp Thr Val Gln Cys Phe Ser Cys His Ala Ala Ile Asp Arg Trp
35 40 45
Gln Tyr Gly Asp Ser Ala Val Gly Arg His Arg Arg Ile Ser Pro Asn
50 55 60
Cys Arg Phe Ile
65

<210> 17

<211> 68
 <212> PRT
 <213> Homo sapiens

<400> 17
 Glu Phe Asn Arg Leu Lys Thr Phe Ala Asn Phe Pro Ser Gly Ser Pro
 1 5 10 15
 Val Ser Ala Ser Thr Leu Ala Arg Ala Gly Phe Leu Tyr Thr Gly Glu
 20 25 30
 Gly Asp Thr Val Arg Cys Phe Ser Cys His Ala Ala Val Asp Arg Trp
 35 40 45
 Gln Tyr Gly Asp Ser Ala Val Gly Arg His Arg Lys Val Ser Pro Asn
 50 55 60
 Cys Arg Phe Ile
 65

<210> 18
 <211> 68
 <212> PRT
 <213> Homo sapiens

<400> 18
 Glu Leu Tyr Arg Met Ser Thr Tyr Ser Thr Phe Pro Ala Gly Val Pro
 1 5 10 15
 Val Ser Glu Arg Ser Leu Ala Arg Ala Gly Phe Tyr Tyr Thr Gly Val
 20 25 30
 Asn Asp Lys Val Lys Cys Phe Cys Cys Gly Leu Met Leu Asp Asn Trp
 35 40 45
 Lys Arg Gly Asp Ser Pro Thr Glu Lys His Lys Lys Leu Tyr Pro Ser
 50 55 60
 Cys Arg Phe Val
 65

<210> 19
 <211> 68
 <212> PRT
 <213> Homo sapiens

<400> 19
 Glu Leu Tyr Arg Met Ser Thr Tyr Ser Thr Phe Pro Ala Gly Val Pro
 1 5 10 15
 Val Ser Glu Arg Ser Leu Ala Arg Ala Gly Phe Tyr Tyr Thr Gly Val
 20 25 30
 Asn Asp Lys Val Lys Cys Phe Cys Cys Gly Leu Met Leu Asp Asn Trp
 35 40 45
 Lys Leu Gly Asp Ser Pro Ile Gln Lys His Lys Gln Leu Tyr Pro Ser
 50 55 60
 Cys Ser Phe Ile
 65

<210> 20
 <211> 68
 <212> PRT
 <213> Mus musculus

<400> 20

Glu	Glu	Ala	Arg	Leu	Lys	Ser	Phe	Gln	Asn	Trp	Pro	Asp	Tyr	Ala	His
1				5					10					15	
Leu	Thr	Pro	Arg	Glu	Leu	Ala	Ser	Ala	Gly	Leu	Tyr	Tyr	Thr	Gly	Ala
			20					25					30		
Asp	Asp	Gln	Val	Gln	Cys	Phe	Cys	Cys	Gly	Gly	Lys	Leu	Lys	Asn	Trp
		35					40					45			
Glu	Pro	Cys	Asp	Arg	Ala	Trp	Ser	Glu	His	Arg	Arg	His	Phe	Pro	Asn
	50					55					60				
Cys	Phe	Phe	Val												
65															

<210> 21
 <211> 68
 <212> PRT
 <213> Homo sapiens

Glu	Glu	Ala	Arg	Leu	Lys	Ser	Phe	Gln	Asn	Trp	Pro	Asp	Tyr	Ala	His
1				5					10					15	
Leu	Thr	Pro	Arg	Glu	Leu	Ala	Ser	Ala	Gly	Leu	Tyr	Tyr	Thr	Gly	Ile
			20					25					30		
Gly	Asp	Gln	Val	Gln	Cys	Phe	Cys	Cys	Gly	Gly	Lys	Leu	Lys	Asn	Trp
		35					40					45			
Glu	Pro	Cys	Asp	Arg	Ala	Trp	Ser	Glu	His	Arg	Arg	His	Phe	Pro	Asn
	50					55					60				
Cys	Phe	Phe	Val												
65															

<210> 22
 <211> 67
 <212> PRT
 <213> Homo sapiens

Glu	Asn	Ala	Arg	Leu	Leu	Thr	Phe	Gln	Thr	Trp	Pro	Leu	Thr	Phe	Leu
1				5					10					15	
Ser	Pro	Thr	Asp	Leu	Ala	Arg	Ala	Gly	Phe	Tyr	Tyr	Ile	Gly	Pro	Gly
			20					25					30		
Asp	Arg	Val	Ala	Cys	Phe	Ala	Cys	Gly	Gly	Lys	Leu	Ser	Asn	Trp	Glu
		35					40					45			
Pro	Lys	Asp	Asn	Ala	Met	Ser	Glu	His	Leu	Arg	His	Phe	Pro	Lys	Cys
	50					55					60				
Pro	Phe	Ile													
65															

<210> 23
 <211> 67
 <212> PRT
 <213> Homo sapiens

Glu	Glu	Ala	Arg	Phe	Leu	Thr	Tyr	His	Met	Trp	Pro	Leu	Thr	Phe	Leu
1				5					10					15	
Ser	Pro	Ser	Glu	Leu	Ala	Arg	Ala	Gly	Phe	Tyr	Tyr	Ile	Gly	Pro	Gly
			20					25					30		
Asp	Arg	Val	Ala	Cys	Phe	Ala	Cys	Gly	Gly	Lys	Leu	Ser	Asn	Trp	Glu

<210> 27
 <211> 68
 <212> PRT
 <213> Homo sapiens

<400> 27
 His Ala Ala Arg Met Arg Thr Phe Met Tyr Trp Pro Ser Ser Val Pro
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 Val Gln Pro Glu Gln Leu Ala Ser Ala Gly Phe Tyr Tyr Val Gly Arg
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 Asn Asp Asp Val Lys Cys Phe Gly Cys Asp Gly Gly Leu Arg Cys Trp
 35 40 45
 Glu Ser Gly Asp Asp Pro Trp Val Glu His Ala Lys Trp Phe Pro Arg
 50 55 60
 Cys Glu Phe Leu
 65

<210> 28
 <211> 68
 <212> PRT
 <213> Orgyia pseudotsugata

<400> 28
 Glu Ala Ala Arg Leu Arg Thr Phe Ala Glu Trp Pro Arg Gly Leu Lys
 1 5 10 15
 Gln Arg Pro Glu Glu Leu Ala Glu Ala Gly Phe Phe Tyr Thr Gly Gln
 20 25 30
 Gly Asp Lys Thr Arg Cys Phe Cys Cys Asp Gly Gly Leu Lys Asp Trp
 35 40 45
 Glu Pro Asp Asp Ala Pro Trp Gln Gln His Ala Arg Trp Tyr Asp Arg
 50 55 60
 Cys Glu Tyr Val
 65

<210> 29
 <211> 68
 <212> PRT
 <213> Cydia pomonella

<400> 29
 Glu Ala Ala Arg Val Lys Ser Phe His Asn Trp Pro Arg Cys Met Lys
 1 5 10 15
 Gln Arg Pro Glu Gln Met Ala Asp Ala Gly Phe Phe Tyr Thr Gly Tyr
 20 25 30
 Gly Asp Asn Thr Lys Cys Phe Tyr Cys Asp Gly Gly Leu Lys Asp Trp
 35 40 45
 Glu Pro Glu Asp Val Pro Trp Glu Gln His Val Arg Trp Phe Asp Arg
 50 55 60
 Cys Ala Tyr Val
 65

<210> 30
 <211> 68
 <212> PRT
 <213> Drosophila melanogaster

<400> 30

Val Asp Ala Arg Leu Arg Thr Phe Thr Asp Trp Pro Ile Ser Asn Ile
1 5 10 15
Gln Pro Ala Ser Ala Leu Ala Gln Ala Gly Leu Tyr Tyr Gln Lys Ile
20 25 30
Gly Asp Gln Val Arg Cys Phe His Cys Asn Ile Gly Leu Arg Ser Trp
35 40 45
Gln Lys Glu Asp Glu Pro Trp Phe Glu His Ala Lys Trp Ser Pro Lys
50 55 60
Cys Gln Phe Val
65

<210> 31

<211> 66

<212> PRT

<213> *Drosophila melanogaster*

<400> 31

Glu Ser Val Arg Leu Ala Thr Phe Gly Glu Trp Pro Leu Asn Ala Pro
1 5 10 15
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20 25 30
Glu Ala Glu Cys Asp Phe Cys His Val Arg Ile Asp Arg Trp Glu Tyr
35 40 45
Gly Asp Leu Val Ala Glu Arg His Arg Arg Ser Ser Pro Ile Cys Ser
50 55 60
Met Val
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<210> 32

<211> 46

<212> PRT

<213> *Homo sapiens*

<400> 32

Glu Gln Leu Arg Arg Leu Gln Glu Glu Arg Thr Cys Lys Val Cys Met
1 5 10 15
Asp Lys Glu Val Ser Val Val Phe Ile Pro Cys Gly His Leu Val Val
20 25 30
Cys Gln Glu Cys Ala Pro Ser Leu Arg Lys Cys Pro Ile Cys
35 40 45

<210> 33

<211> 46

<212> PRT

<213> *Homo sapiens*

<400> 33

Glu Gln Leu Arg Arg Leu Pro Glu Glu Arg Thr Cys Lys Val Cys Met
1 5 10 15
Asp Lys Glu Val Ser Ile Val Phe Ile Pro Cys Gly His Leu Val Val
20 25 30
Cys Lys Asp Cys Ala Pro Ser Leu Arg Lys Cys Pro Ile Cys
35 40 45

<210> 34
 <211> 46
 <212> PRT
 <213> Mus musculus

<400> 34
 Glu Gln Leu Arg Arg Leu Gln Glu Glu Lys Leu Ser Lys Ile Cys Met
 1 5 10 15
 Asp Arg Asn Ile Ala Ile Val Phe Phe Pro Cys Gly His Leu Ala Thr
 20 25 30
 Cys Lys Gln Cys Ala Glu Ala Val Asp Lys Cys Pro Met Cys
 35 40 45

<210> 35
 <211> 46
 <212> PRT
 <213> Homo sapiens

<400> 35
 Glu Gln Leu Arg Arg Leu Gln Glu Glu Lys Leu Cys Lys Ile Cys Met
 1 5 10 15
 Asp Arg Asn Ile Ala Ile Val Phe Val Pro Cys Gly His Leu Val Thr
 20 25 30
 Cys Lys Gln Cys Ala Glu Ala Val Asp Lys Cys Pro Met Cys
 35 40 45

<210> 36
 <211> 46
 <212> PRT
 <213> Drosophila melanogaster

<400> 36
 Glu Glu Asn Arg Gln Leu Lys Asp Ala Arg Leu Cys Lys Val Cys Leu
 1 5 10 15
 Asp Glu Glu Val Gly Val Val Phe Leu Pro Cys Gly His Leu Ala Thr
 20 25 30
 Cys Asn Gln Cys Ala Pro Ser Val Ala Asn Cys Pro Met Cys
 35 40 45

<210> 37
 <211> 46
 <212> PRT
 <213> Cydia pomonella

<400> 37
 Glu Lys Glu Pro Gln Val Glu Asp Ser Lys Leu Cys Lys Ile Cys Tyr
 1 5 10 15
 Val Glu Glu Cys Ile Val Cys Phe Val Pro Cys Gly His Val Val Ala
 20 25 30
 Cys Ala Lys Cys Ala Leu Ser Val Asp Lys Cys Pro Met Cys
 35 40 45

<210> 38
 <211> 46
 <212> PRT

<213> *Orgyia pseudotsugata*

<400> 38

Ala	Val	Glu	Ala	Glu	Val	Ala	Asp	Asp	Arg	Leu	Cys	Lys	Ile	Cys	Leu
1				5					10					15	
Gly	Ala	Glu	Lys	Thr	Val	Cys	Phe	Val	Pro	Cys	Gly	His	Val	Val	Ala
			20					25					30		
Cys	Gly	Lys	Cys	Ala	Ala	Gly	Val	Thr	Thr	Cys	Pro	Val	Cys		
		35					40					45			

<210> 39

<211> 2474

<212> DNA

<213> *Mus musculus*

<400> 39

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ttttcctgtg	agctgtaccg	attgtccacg	tattcagctt	ttcccagggg	agttcctgtg	300
tcagaaagga	gtctggctcg	tgctggcttt	tactacactg	gtgccaatga	caaggtcaag	360
tgcttctgct	gtggcctgat	gctagacaac	tggaacaacg	gggacagtcc	catggagaag	420
cacagaaaat	tgtaccccag	ctgcaacttt	gtacagactt	tgaatccagc	caacagtctg	480
gaagctagtc	ctcggccttc	tcttccttcc	acggcgatga	gcacccatgcc	tttgagcttt	540
gcaagtcttg	agaatactgg	ctatttcagt	ggctcttact	cgagctttcc	ctcagaccct	600
gtgaacttcc	gagcaaatca	agattgtcct	gctttgagca	caagtcccta	ccactttgca	660
atgaacacag	agaaggccag	attactcacc	tatgaaacat	ggccattgtc	ttttctgtca	720
ccagcaaagc	tgcccaaagc	aggcttctac	tacataggac	ctggagatag	agtggcctgc	780
tttgctgctg	atgggaaact	gagcaactgg	gaacgtaagg	atgatgctat	gtcagagcac	840
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actgtctcta	acctgagcat	gcagacacac	gcagcccgtg	ttagaacatt	ctctaactgg	960
ccttctagtg	cactagttca	ttcccaggaa	cttgcaagtg	cgggctttta	ttatacagga	1020
cacagtgatg	atgtcaagtg	tttatgctgt	gatggtgggc	tgaggtgctg	ggaatctgga	1080
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ctaattggtcc	atggttgcaa	cttcagccag	gaggaagtgc	actgtcactc	ccagttccat	2040
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gaaaaacttt	tgtctgaagt	caagaatgaa	tgaattactt	atataataat	tttaattggt	2160
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ggctagaatc	catgaaccaa	gctgcaaaga	tctcacgcta	aataaggcgg	aaagatttgg	2400
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<210> 40
 <211> 602
 <212> PRT
 <213> Mus musculus

<400> 40

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Ala	Asp	Thr	Phe	Glu	Leu	Lys	Tyr	Asp	Phe	Ser	Cys	Glu	Leu	Tyr	Arg
			20					25					30		
Leu	Ser	Thr	Tyr	Ser	Ala	Phe	Pro	Arg	Gly	Val	Pro	Val	Ser	Glu	Arg
		35					40					45			
Ser	Leu	Ala	Arg	Ala	Gly	Phe	Tyr	Tyr	Thr	Gly	Ala	Asn	Asp	Lys	Val
	50					55					60				
Lys	Cys	Phe	Cys	Cys	Gly	Leu	Met	Leu	Asp	Asn	Trp	Lys	Gln	Gly	Asp
65					70					75				80	
Ser	Pro	Met	Glu	Lys	His	Arg	Lys	Leu	Tyr	Pro	Ser	Cys	Asn	Phe	Val
				85					90					95	
Gln	Thr	Leu	Asn	Pro	Ala	Asn	Ser	Leu	Glu	Ala	Ser	Pro	Arg	Pro	Ser
			100					105					110		
Leu	Pro	Ser	Thr	Ala	Met	Ser	Thr	Met	Pro	Leu	Ser	Phe	Ala	Ser	Ser
		115					120					125			
Glu	Asn	Thr	Gly	Tyr	Phe	Ser	Gly	Ser	Tyr	Ser	Ser	Phe	Pro	Ser	Asp
	130					135					140				
Pro	Val	Asn	Phe	Arg	Ala	Asn	Gln	Asp	Cys	Pro	Ala	Leu	Ser	Thr	Ser
145					150					155					160
Pro	Tyr	His	Phe	Ala	Met	Asn	Thr	Glu	Lys	Ala	Arg	Leu	Leu	Thr	Tyr
				165					170					175	
Glu	Thr	Trp	Pro	Leu	Ser	Phe	Leu	Ser	Pro	Ala	Lys	Leu	Ala	Lys	Ala
			180					185					190		
Gly	Phe	Tyr	Tyr	Ile	Gly	Pro	Gly	Asp	Arg	Val	Ala	Cys	Phe	Ala	Cys
		195					200					205			
Asp	Gly	Lys	Leu	Ser	Asn	Trp	Glu	Arg	Lys	Asp	Asp	Ala	Met	Ser	Glu
	210					215					220				
His	Gln	Arg	His	Phe	Pro	Ser	Cys	Pro	Phe	Leu	Lys	Asp	Leu	Gly	Gln
225					230					235					240
Ser	Ala	Ser	Arg	Tyr	Thr	Val	Ser	Asn	Leu	Ser	Met	Gln	Thr	His	Ala
				245					250					255	
Ala	Arg	Ile	Arg	Thr	Phe	Ser	Asn	Trp	Pro	Ser	Ser	Ala	Leu	Val	His
			260					265					270		
Ser	Gln	Glu	Leu	Ala	Ser	Ala	Gly	Phe	Tyr	Tyr	Thr	Gly	His	Ser	Asp
		275					280					285			
Asp	Val	Lys	Cys	Leu	Cys	Cys	Asp	Gly	Gly	Leu	Arg	Cys	Trp	Glu	Ser
	290					295					300				
Gly	Asp	Asp	Pro	Trp	Val	Glu	His	Ala	Lys	Trp	Phe	Pro	Arg	Cys	Glu
305					310					315					320
Tyr	Leu	Leu	Arg	Ile	Lys	Gly	Gln	Glu	Phe	Val	Ser	Gln	Val	Gln	Ala
				325					330					335	
Gly	Tyr	Pro	His	Leu	Leu	Glu	Gln	Leu	Leu	Ser	Thr	Ser	Asp	Ser	Pro
		340						345					350		
Glu	Asp	Glu	Asn	Ala	Asp	Ala	Ala	Ile	Val	His	Phe	Gly	Pro	Gly	Glu
		355					360					365			
Ser	Ser	Glu	Asp	Val	Val	Met	Met	Ser	Thr	Pro	Val	Val	Lys	Ala	Ala
	370					375					380				
Leu	Glu	Met	Gly	Phe	Ser	Arg	Ser	Leu	Val	Arg	Gln	Thr	Val	Gln	Trp
385					390					395					400
Gln	Ile	Leu	Ala	Thr	Gly	Glu	Asn	Tyr	Arg	Thr	Val	Ser	Asp	Leu	Val
				405					410					415	
Ile	Gly	Leu	Leu	Asp	Ala	Glu	Asp	Glu	Met	Arg	Glu	Glu	Gln	Met	Glu

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<210> 41
<211> 2416
<212> DNA
<213> Mus musculus
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-23-

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tattgtgttc	attccgtgtg	gtcatctagt	agtctgccag	gaatgtgccc	cttctctaag	1860
gaagtgtccc	atctgcagg	ggacaatcaa	ggggactgtg	cgcacatttc	tctcatgagt	1920
gaagaatggt	ctgaaagtat	tgttggacat	cagaagctgt	cagaacaaag	aatgaactac	1980
tgatttcagc	tcttcagcag	gacattctac	tctctttcaa	gattagtaat	cttgctttat	2040
gaagggtagc	attgtatat	taagcttagt	ctgttgcaag	ggaaggtcta	tgctgttgag	2100
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cttcttggga	tttgggaatt	tggggaaaagc	tttggaatcc	agtgatgtgg	agctcagaaa	2220
tcctggaacc	agtgactctg	gtactcagta	gatagggtac	cctgtacttc	ttgggtgcttt	2280
tccagtctgg	gaaataagga	ggaatctgct	gctggtaaaa	atttgctgga	tgtgagaaat	2340
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<210> 42

<211> 591

<212> PRT

<213> Mus musculus

<400> 42

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			20					25					30		
Ser	Ala	Phe	Pro	Arg	Gly	Val	Pro	Val	Ser	Glu	Arg	Ser	Leu	Ala	Arg
		35					40					45			
Ala	Gly	Phe	Tyr	Tyr	Thr	Gly	Val	Asn	Asp	Lys	Val	Lys	Cys	Phe	Cys
	50					55					60				
Cys	Gly	Leu	Met	Leu	Asp	Asn	Trp	Lys	Gln	Gly	Asp	Ser	Pro	Val	Glu
65					70					75					80
Lys	His	Arg	Gln	Phe	Tyr	Pro	Ser	Cys	Ser	Phe	Val	Gln	Thr	Leu	Leu
			85						90					95	
Ser	Ala	Ser	Leu	Gln	Ser	Pro	Ser	Lys	Asn	Met	Ser	Pro	Val	Lys	Ser
			100					105					110		
Arg	Phe	Ala	His	Ser	Ser	Pro	Leu	Glu	Arg	Gly	Gly	Ile	His	Ser	Asn
		115					120					125			
Leu	Cys	Ser	Ser	Pro	Leu	Asn	Ser	Arg	Ala	Val	Glu	Asp	Phe	Ser	Ser
	130					135					140				
Arg	Met	Asp	Pro	Cys	Ser	Tyr	Ala	Met	Ser	Thr	Glu	Glu	Ala	Arg	Phe
145					150					155					160
Leu	Thr	Tyr	Ser	Met	Trp	Pro	Leu	Ser	Phe	Leu	Ser	Pro	Ala	Glu	Leu
			165						170					175	
Ala	Arg	Ala	Gly	Phe	Tyr	Tyr	Ile	Gly	Pro	Gly	Asp	Arg	Val	Ala	Cys
		180						185					190		
Phe	Ala	Cys	Gly	Gly	Lys	Leu	Ser	Asn	Trp	Glu	Pro	Lys	Asp	Tyr	Ala
	195						200					205			
Met	Ser	Glu	His	Arg	Arg	His	Phe	Pro	His	Cys	Pro	Phe	Leu	Glu	Asn
	210					215					220				
Thr	Ser	Glu	Thr	Gln	Arg	Phe	Ser	Ile	Ser	Asn	Leu	Ser	Met	Gln	Thr
225				230						235					240
His	Ser	Ala	Arg	Leu	Arg	Thr	Phe	Leu	Tyr	Trp	Pro	Pro	Ser	Val	Pro
			245						250					255	
Val	Gln	Pro	Glu	Gln	Leu	Ala	Ser	Ala	Gly	Phe	Tyr	Tyr	Val	Asp	Arg
		260						265					270		
Asn	Asp	Asp	Val	Lys	Cys	Leu	Cys	Cys	Asp	Gly	Gly	Leu	Arg	Cys	Trp
	275						280					285			
Glu	Pro	Gly	Asp	Asp	Pro	Trp	Ile	Glu	His	Ala	Lys	Trp	Phe	Pro	Arg
	290					295					300				
Cys	Glu	Phe	Leu	Ile	Arg	Met	Lys	Gly	Gln	Glu	Phe	Val	Asp	Glu	Ile

305					310					315					320
Gln	Ala	Arg	Tyr	Pro	His	Leu	Leu	Glu	Gln	Leu	Leu	Ser	Thr	Ser	Asp
				325					330					335	
Thr	Pro	Gly	Glu	Glu	Asn	Ala	Asp	Pro	Thr	Glu	Thr	Val	Val	His	Phe
			340					345					350		
Gly	Pro	Gly	Glu	Ser	Ser	Lys	Asp	Val	Val	Met	Met	Ser	Thr	Pro	Val
		355					360					365			
Val	Lys	Ala	Ala	Leu	Glu	Met	Gly	Phe	Ser	Arg	Ser	Leu	Val	Arg	Gln
	370					375				380					
Thr	Val	Gln	Arg	Gln	Ile	Leu	Ala	Thr	Gly	Glu	Asn	Tyr	Arg	Thr	Val
385					390					395					400
Asn	Asp	Ile	Val	Ser	Val	Leu	Leu	Asn	Ala	Glu	Asp	Glu	Arg	Arg	Glu
				405					410					415	
Glu	Glu	Lys	Glu	Arg	Gln	Thr	Glu	Glu	Met	Ala	Ser	Gly	Asp	Leu	Ser
			420					425					430		
Leu	Ile	Arg	Lys	Asn	Arg	Met	Ala	Leu	Phe	Gln	Gln	Leu	Thr	His	Val
	435						440					445			
Leu	Pro	Ile	Leu	Asp	Asn	Leu	Glu	Ala	Ser	Val	Ile	Thr	Lys	Gln	
	450					455				460					
Glu	His	Asp	Ile	Ile	Arg	Gln	Lys	Thr	Gln	Ile	Pro	Leu	Gln	Ala	Arg
465					470					475					480
Glu	Leu	Ile	Asp	Thr	Val	Leu	Val	Lys	Gly	Asn	Ala	Ala	Ala	Asn	Ile
				485					490					495	
Phe	Lys	Asn	Ser	Leu	Lys	Gly	Ile	Asp	Ser	Thr	Leu	Tyr	Glu	Asn	Leu
		500						505					510		
Phe	Val	Glu	Lys	Asn	Met	Lys	Tyr	Ile	Pro	Thr	Glu	Asp	Val	Ser	Gly
		515					520					525			
Leu	Ser	Leu	Glu	Glu	Gln	Leu	Arg	Arg	Leu	Gln	Glu	Glu	Arg	Thr	Cys
	530					535					540				
Lys	Val	Cys	Met	Asp	Arg	Glu	Val	Ser	Ile	Val	Phe	Ile	Pro	Cys	Gly
545					550					555					560
His	Leu	Val	Val	Cys	Gln	Glu	Cys	Ala	Pro	Ser	Leu	Arg	Lys	Cys	Pro
				565					570					575	
Ile	Cys	Arg	Gly	Thr	Ile	Lys	Gly	Thr	Val	Arg	Thr	Phe	Leu	Ser	
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<210> 43
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetic based on viral sequence

<400> 43
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<210> 44
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic primer based on Homo sapiens

<400> 44

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21

<210> 45

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic primer based on Homo sapiens

<400> 45

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25